



CSIR-NET

Council of Scientific & Industrial Research

LIFE SCIENCE

VOLUME – 6

**DEVELOPMENT BIOLOGY, PLANT
& ANIMAL PHYSIOLOGY**



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GENETICS (INHERITANCE BIOLOGY)

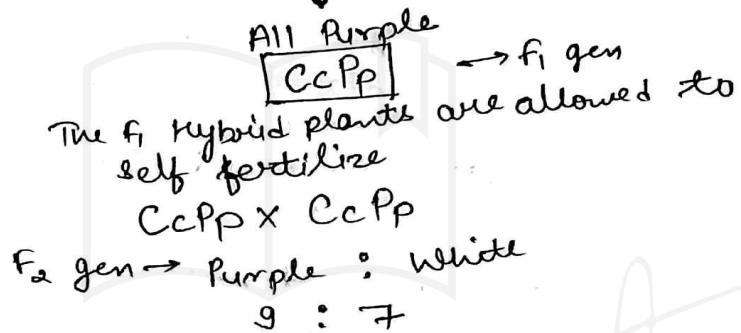
Gene interaction :-

→ When two or more different genes influence the outcome of single trait, this is known as a gene interaction.

Discovery - William Bateson & Reginald Punnett in 1906. They discovered an unexpected gene interaction when they studied crosses involving the sweet pea, *Lathyrus odoratus*.

ex. P generation white flowered plant \times white flowered plant
 $CCPP \quad ccPP$

9:3:3:1
 ↓ convert
 9:7



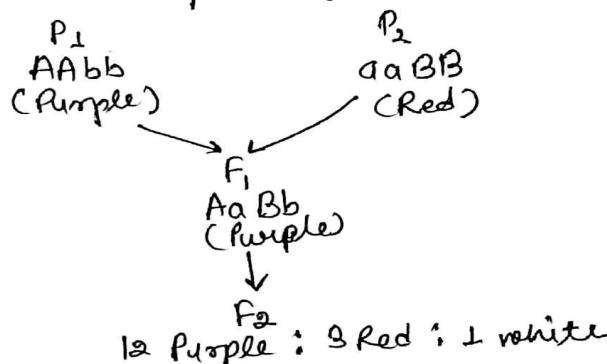
Epistasis :-

Epistasis → Greek for standing upon
 A Type of gene interaction where one gene masks or modifies the expression of another gene at distinct loci.
 → Any gene that masks the expression of another non-allelic gene is epistatic to that gene.
 → The gene suppressed is hypostatic.
 → Epistasis is diff from dominance. Epistasis is the interaction b/w diff genes (non-alleles). Dominance is the interaction b/w diff alleles of the same gene that is intra allelic.

① Dominant Epistasis :-

Dominant allele of one gene masks the effects of either allele of the second gene. It is termed as dominant epistasis

ex:-



* Recessive epistasis:-

In the case of recessive epistasis, in a pair of non-allelic genes one produces its phenotypic effect independently in a dominant state, but another cannot produce a phenotypic effect independent.

Ratio 9:3:3:1 become 9:3:4

Purple : Red : White

* Duplicate recessive epistasis:- (complementary gene interaction)

If two non-allelic genes are involved in a specific pathway & functional products from both are required for expression, then one homozygous recessive allele in either allele pair would result in the mutant phenotype.

9:3:3:1 → 9:7

* Duplicate dominant interaction:-

9:3:3:1 → 15:1

* Pleiotropy:-

→ Most of the biochemical pathways in the living organism are interconnected.

→ The term pleiotropy refers to the effect of a single gene on more than one character/ trait.

Some time one trait will be very evident & others will be less evident.

- Several characteristics may differ b/w individuals belonging to the same spp. These differences are termed variations.
- The mechanism of transmission of characters, resemblances as well as differences, from the parental generation to the offspring is heredity. (~~is different~~)
- The study of heredity, variations & env. factor known as genetics.

Greek word genno = give birth

Genetics → (S + greeks)

① Classical genetics → Mendel's principle, Sex determination

② Molecular genetics → Genetic material

③ Evolutionary genetics → Popn genetics

Gregor Johann Mendel (1822-1884)

Rediscovered the Mendel's law - ① Hugo de Vries (Holland)
 (1900) ② Carl Correns (Germany)
 ③ Erich Tschermak (Austria)

- ⇒ He discovered that individual traits are inherited as discrete factors which retain their physical identity in a hybrid.
- ⇒ Later, these factors come to be known as gene.
- ⇒ A gene is defined as a unit of heredity.

• Allele :-

Each gene may exists in alternative forms known as allele.

• Homologous - chromosome that carry the same set of genes in the same seq., although they may not necessarily carry identical alleles of each gene

* Rh Blood group System :-

- Rh antigen & Factor tut on the plasma membrane.
- Marcus rhesus → monkey
- The Rh antigens are located on two Rhesus protein RhD and RHCE
- RH D and RHCE are expressed only in RBCs.
- More than 170 alleles of RH D gene have been found.
- Rh D protein carries for D antigen
- Rh CE protein carries for CE antigen in various combination (Cc, Cc, cE or CE)
- The classification of Rh-positive and Rh-negative individuals is determined by the presence or absence of the D antigen on the surface of RBCs.
- If absent Rh⁻ D antigen tut on a person's RBC, the person is Rh⁺
- if absent Rh⁻ Erythroblastosis fetalis (or hemolytic disease)

* Lethal allele :-

The alleles created by mutation in these genes are called lethal alleles. Lethal alleles may be recessive or dominant.

4 categories

- ① Early onset → early death of an organism
- ② Late onset → delayed effect
- ③ Conditional → kill organism under certain env. conditions
- ④ Semilethal → kill only some individuals but not all.

* Penetrance :-

The percentage of individuals that shows a particular phenotype among those capable of showing it, is known as penetrance. ex. - Polydactyly in humans.

(More than 5 fingers)

- A particular gene may produce diff. degrees of expression in diff. individuals. This is known as expressivity.

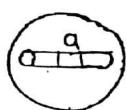
* Phenocopy :-

A phenotype that is not genetically controlled but looks like a genetically controlled one is cl phenocopy. It is an environmentally induced phenotype that resembles the phenotype determined by the genotype.

ex. → phenocopy of vit-D resistant rickets.

LAW OF GENETICS

- Allele:- Alternate form of the gene
 - Different nose shape - Polymorphism
 - Eye colour is A gene but diff. Eye colour like blue, black, white eye is Allele
 - Mutation creates Allele.
 - Local population in a particular area known as wild type if it is present in maximum popn
- Diff. nose shape
- | | | |
|------------|------------|------------|
| <u>AGT</u> | <u>A@T</u> | <u>ATT</u> |
| | ↓ | |
| mutation | | |
- Haploid - Diploid Tetraploid



Diploid



Tetraploid



→ Alleles present on chromosome different allele representation.

Eye colour - (Black eye - b^+ , B, B^1 , B^A)

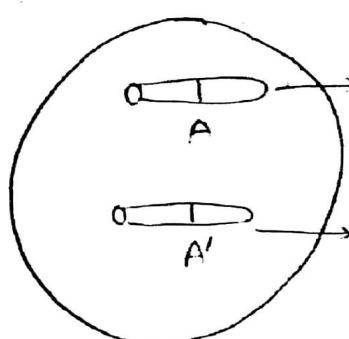
* Diploid - two alleles of a gene

Haploid - Single alleles of a gene

→ Crene is not seen → The wrinkled phenotype of seed is due to the absence of Amylo pectin

* Law of Genetics

① Law of Dominance:- → Dominant & Recessive are not the properties of alleles in gene. It is a relationship b/w 2 alleles, that is observed (Recessive) in Phenotype.



Tc, Ts
low

→ β (White → Black)

Protein, UPS ↑, life ↓ (Protein degradation)



Tc, Ts ↑
(Dominant)

Protein ↑ β → $\text{ICat T, life } \uparrow$

Turn over rate ↑

Km low

Affinity ↑

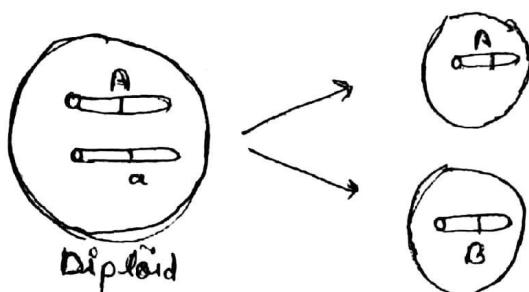
(10^8 -1 sec bind)

UPS ↓

→ Blue colour is shown in ~~next~~ generation.

White → Blue

② Law of Segregation :- (Law of purity of gamete)



① Law of Dominance :-

→ One allele of a gene can cancel the effect of other allele of same gene

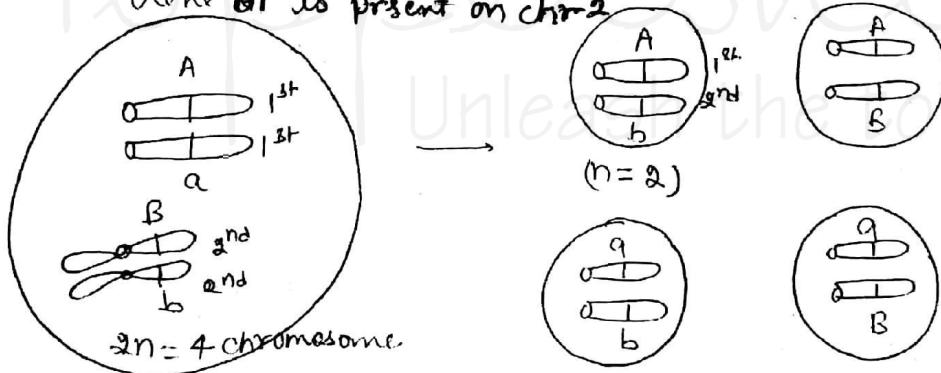
③ Law of Segregation :-

Gametes are Haploid

③ Law of Independent assortment :-

Gene **A** is present on Chr-1

Gene **B** is present on Chr-2



Phenotype = 1:1:1:1

→ Allele can combine with any other allele of other gene
two genes are not on diff chromosome

→ One allele of a gene can combine with any other allele of other gene. This is known as law of Independent assortment.

→ When 2 genes are not on diff chromosome their gametic ratio is 1:1:1:1

LINKAGE

Linkage :-

- two or more genes reside on the same chromosome, they are said to be linked and their transmission pattern is called linkage.
- They may be linked together on the autosomes or on the sex chromosome. Gene that on diff. non-homologous chromosomes are called Unlinked gene.
- linked gene (genes on the same chromosome), however tends to stay together during the formation of gametes.
 linked gene - on same chromosome
 Unlinked gene - (on different chromosome)

* Limits of recombination :-

If two gene loci are, so far, apart in the chromosome that the probability of a chiasma forming b/w them is 100%, then 50% of the gametes will be the parental type (non-crossover) & 50% recombinant (crossover) type. When such dihybrid individuals are testcrossed, they are expected to produce progeny in a 1: 1: 1: 1 ratio as would be expected for genes on diff. chromosome. Recombination b/w two linked genes cannot exceed 50%, even when multiple crossovers occurs b/w them.

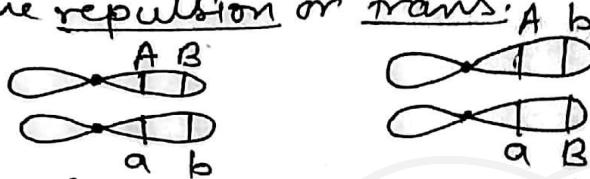
* Morgan's work :-

The effects of linkage were first evident in the result of a dihybrid cross in sweet pea by Bateson & Punnett in 1906.

- Morgan who first discovered the phenomenon of linkage. Morgan postulated that separation of linked gene occurs due to form of chiasmata which represent points of genetic exchange. He used the term crossing over to describe the physical exchange leading to recombination.
- Morgan also proposed that two genes located relatively close to each other along a chromosome are less likely to have a chiasma b/w them as compared to the two genes which are farther apart on the chromosome.
- Drosophila is unusual in that crossing over (hence recombination) does not occur in male. The results is that alleles located in a particular chromosomes show complete linkage in drosophila males. Crossing over is also completely suppressed in female silkworms.

* Cis and trans configuration :-

Doubly heterozygous genotypes can be in two diff. configurations. When two linked genes on each chromosome are of the same type (i.e. both dominant, AB, or both recessive ab) the arrangement is cis & if the coupling is the opposite (i.e. one dominant and one recessive allele, Ab or aB) the arrangement is the repulsion or trans.



Cis (AB/ab)

Trans (Ab/aB)

↓
AB = Parental combination

Ab = Recombinant

Ab = Parental combination

AB = Recombinant.

* Genetic mapping :-

The linkage of the gene in a chromosome can be represented in the form of a genetic map or linkage map or chromosomal map.

Discovery - Alfred Sturtevant

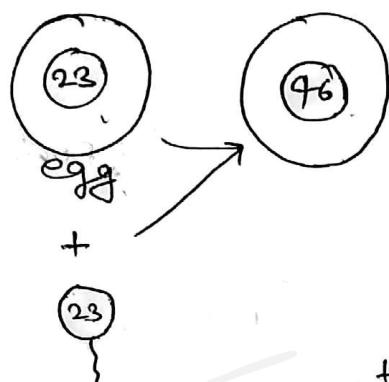
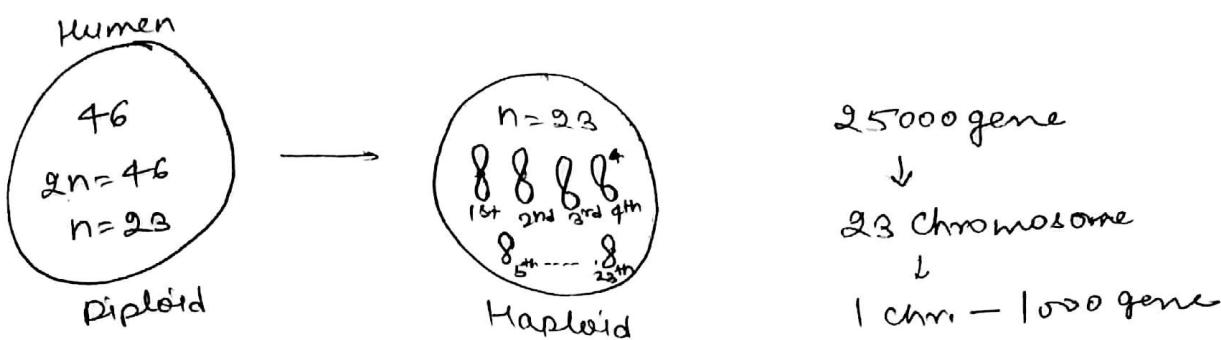
→ He used the frequency of crossing over (i.e. recombination frequency) b/w two genes to prepare the first genetic map of X-chromosome of Drosophila.
1% crossing over or recombination is equivalent to 1 map unit or 1 cM.

By analyzing percent recombination among the progeny of parents that are heterozygous for a no. of linked genes, a genetic map that places the gene in a linear array can be constructed so.

$$\text{Map distance} = \frac{\text{No. of Recombinant offspring}}{\text{Total no. of offspring}} \times 100$$

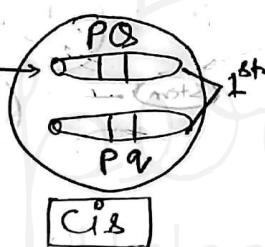
Unit = Map Unit or Centimorgan (cM)

$$1\text{ CM} = 1 \times 10^8 \text{ bp}$$



Linked genes:— When two or more genes ^(Pair of the gene) sit on same chromosome they are known as linked genes

Link gene separated
on crossing over

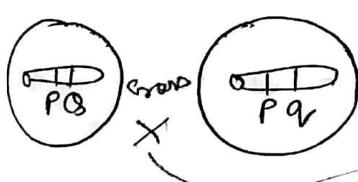


→ Dominant on one chromosome

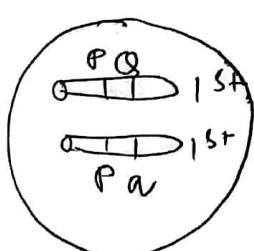
(P Q are gene not allele)
→ Dominant & Recessive both chromosome on 1

Trans

→ One dominant allele & one recessive allele



The linked genes sit on same chromosome
→ Chromosomal map = T. H Morgan

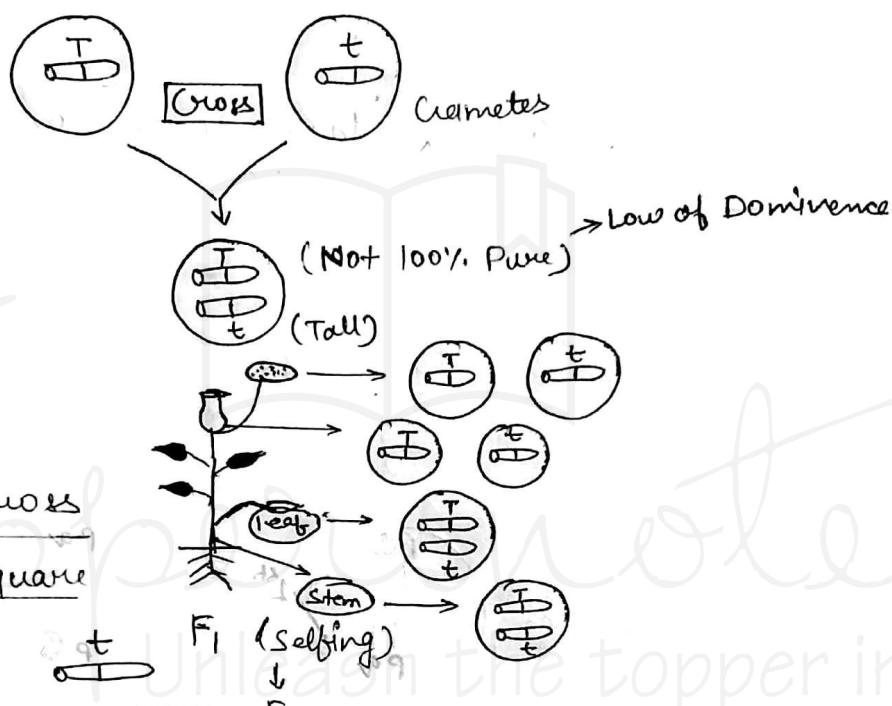
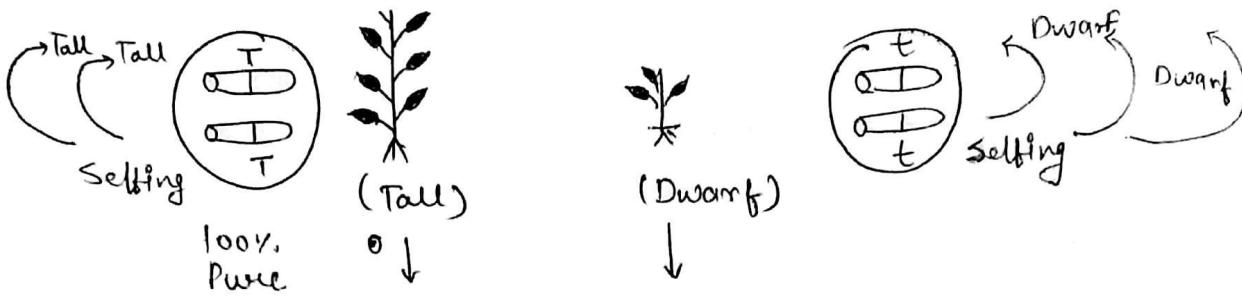


phenotype = 1 : 0 : 0 : 1



Genetic ratio deviates from 1:1:1:1 and it is 1:0:0:1
linkage group = Haploid no. of chromosome are considered as linked group
Haploid no. of chromosome & sex chromosome
Female = 22+X Female = 22+X+Y

MONOHYBRID AND DIHYBRID CROSS



Mono Hybrid cross

↓
Punnett square

♀ T	♂ T	♂ t	
♀ T	Tall	Tall	
♂ t	Tall	Small	Small

F₂ gen all dwarf are pure

Phenotype

3:1

Tall: Small Dwarf

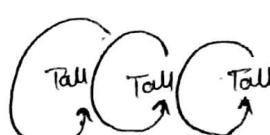
Genotype = 1:2:1

$\frac{1}{4}$, $\frac{2}{4}$, $\frac{1}{4}$

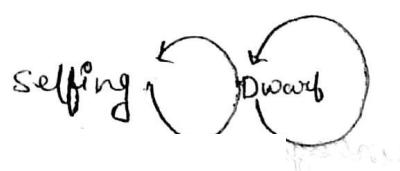
Tall, Tall, Tall,矮

Tt = 3:1

50% pure



TT	Tt
Tall	Tall
Tt	Tt
Tall	Small





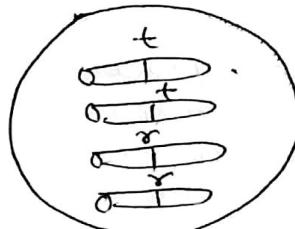
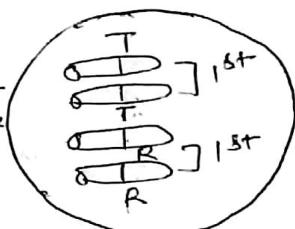
Tall & Round

Dihybrid cross

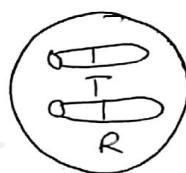


Dwarf & wrinkled

I. A ← Diff Chromosome
↓
1:1:1:1



↓ Gamete for



↓ F1 gen



Dihybrid cross

→

Tall & Round

Tall & Round

Phenotype :-

Tall &
Round
g

: Tall
wrinkled

: Dwarf
Round

: Dwarf
wrinkled
1

Male

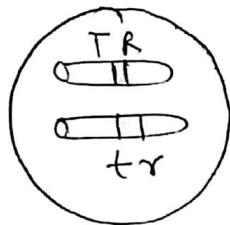
Gamete



Female



linked gene



→ Linked gene में दोनों जिससे Tall & Round, Small & Wrinkled हो सकते हैं और नहीं हो सकते।

Blue eyes
&
Normal Nose

Black
eye
&
Pointed Nose

Gametes -

- ① Blue eyes
&
Normal Nose
- ② Black eyes
&
Normal Nose

- ① Blue eyes
&
Pointed Nose
- ② Black eyes
&
Pointed Nose

Punnett square

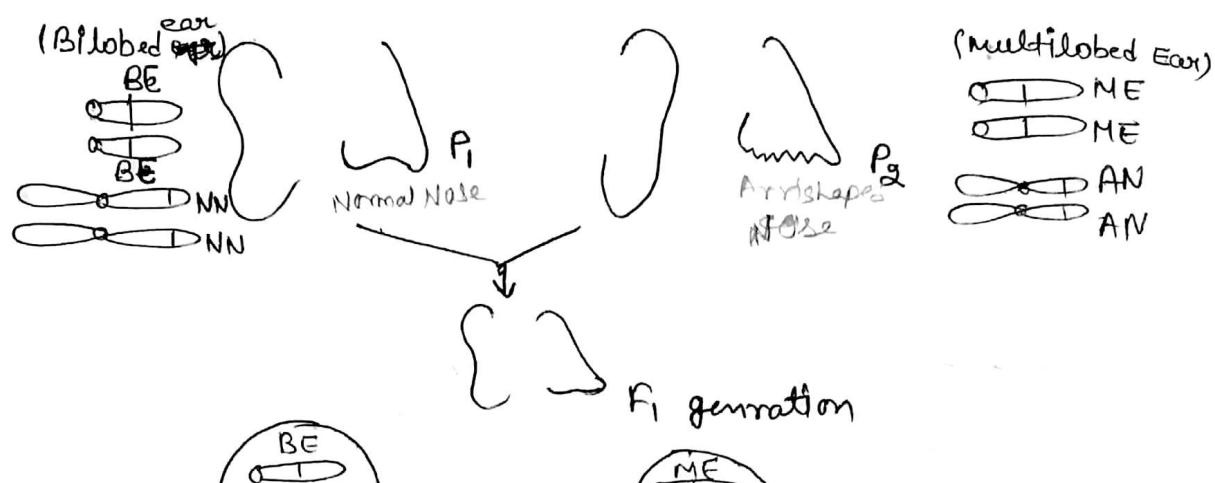
♀	♂	TR	T r	tR	t r
TR	TTRR Tall & Round	TTR r Tall & Round	TtRR Tall & Round	TtR r Tall & Round	TtR r Tall & Round
T r	TTR r Tall & Round	TT r r Tall & Wrinkled	TtR r Tall & Round	TtR r Tall & Round	Tt r r Tall & Wrinkled
tR	TtRR Tall & Round	TtR r Tall & Round	TtRR Small & Round	TtR r Small & Round	ttrr Small & Wrinkled
t r	TtR r Tall & Round	Tt r r Tall & Wrinkled	TtR r Small & Round	Tt r r Small & Round	ttrr Small & Wrinkled

Phenotype :- 9: 3: 3: 1

Genotype :- 1: 2: 2: 4: 1: 2: 1: 2: 1

→ All alleles for any particular gene are found at a specific place on a chromosome called the locus for that gene.

* Test cross - The cross b/w F_1 & its recessive

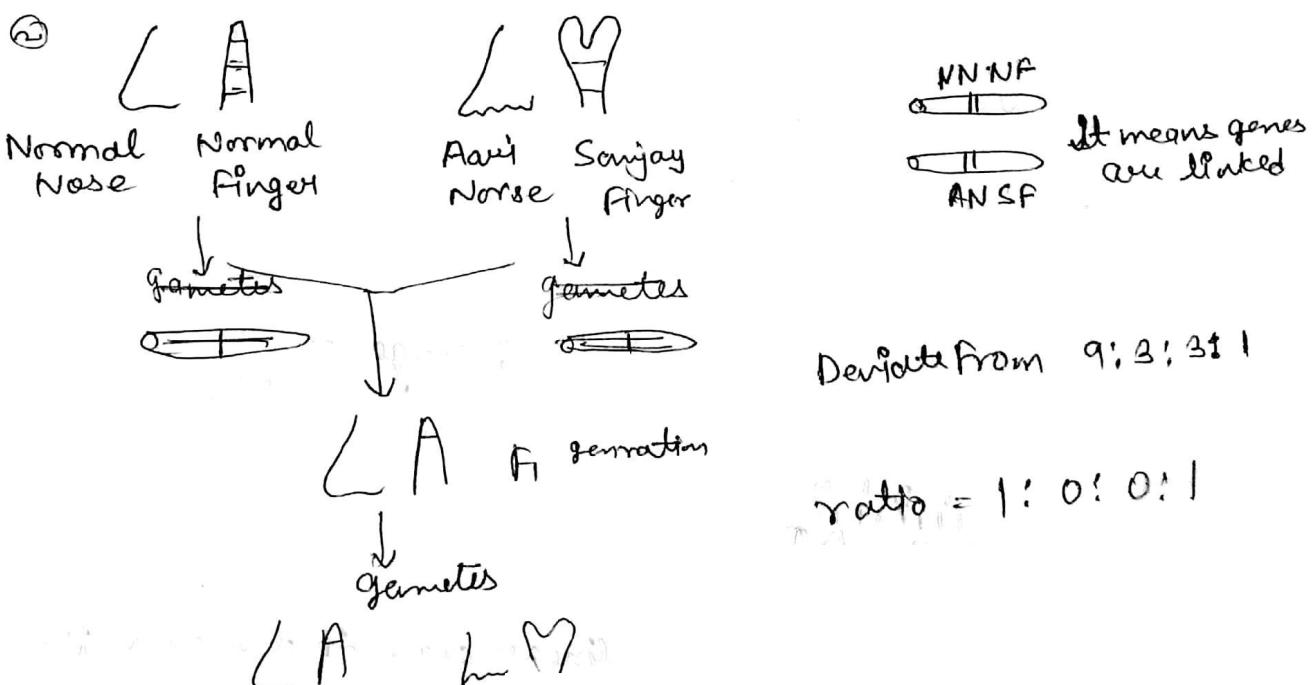


Linkage group :-

→ All the genes that are on a particular chromosome form a linkage group. ex-

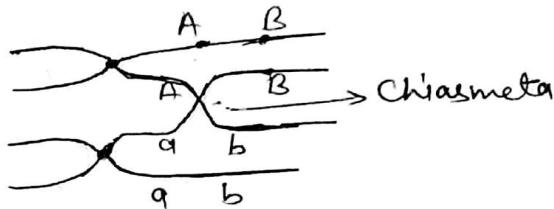
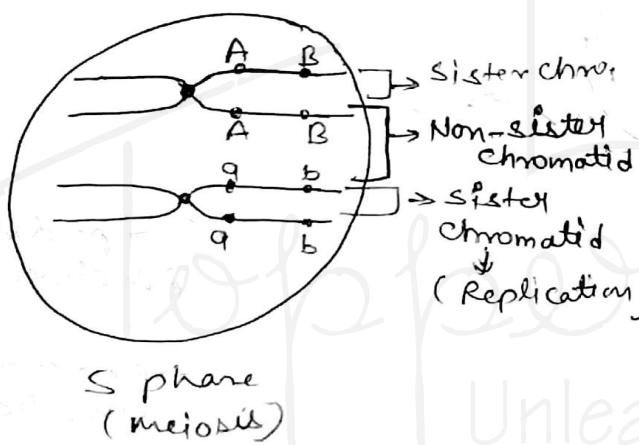
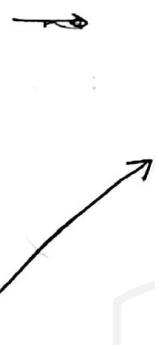
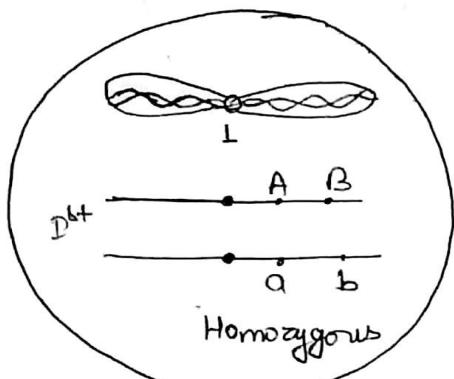
In humans - Male = 22 pairs autosome + 1X chro. + 1Y chro.
= 24 linkage group

Female = 22 pair autosome + 2X chromosome
= 23 linkage group



Crossing over means recombination.

Chiasmeta = Evidence of crossing over



- Normally 1 chiasmeta is fint.
- 4 & 5 very rare
- Chiasmeta no. is variable
- like -

$$A = 1 \quad C = 4$$

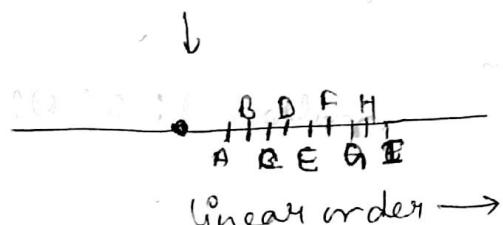
$$B = 3 \quad D = 2$$

$$E = 7 \rightarrow \text{Fasile}$$

→ crossing over fint in non-sister chromatid.

→ In ~~one~~ chromosome on one crossing over second crossing is not found. bcoz. by crossing over chromosome is break

→ 1 chromosome = 1 DNA



$$\text{Linkage d} = \frac{1}{\text{Distance}}$$

→ ~~more~~ distance ~~more~~ ~~less~~ linkage ~~more~~ ~~less~~ 1

→ genes are arranged in linear order that means distance b/w diff genes pair is diff.

Linkage frequency -

$$A-B = 90\%$$

$$A-C = 10\%$$

$$A-D = 20\%$$

$$A-E = 60\%$$

$$A-F = 89\%$$

See \rightarrow A F B E D C

\rightarrow F.A के पास

L \propto 1
crossing over

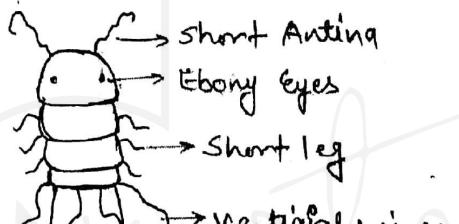
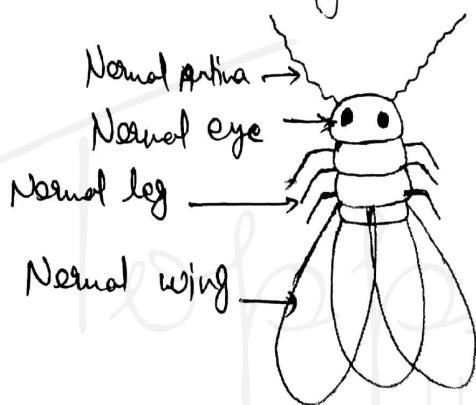
crossing over \uparrow \Rightarrow Linkage FA

\rightarrow Crossing over is a feature of germ cell.

\rightarrow Following type 2 identified दोहरी ये -

① according to F_1 gen. gametes

② By selfing

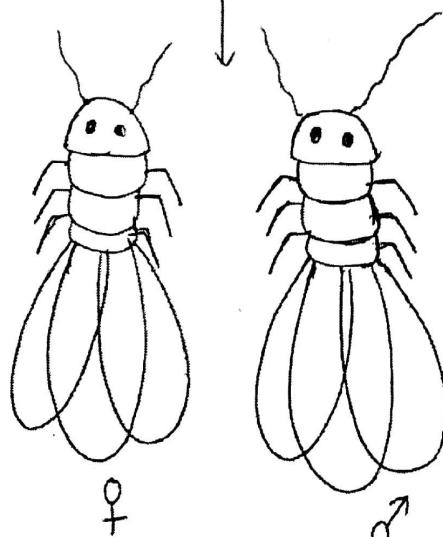


Drosophila

Selfing

$$9 : 3 : 3 : 1$$

N.W ♀	N.W ♂	V.W ♀	V.W. ♂
N.leg	S.leg	N.leg	S.leg



F_1 generation
(Heterozygous)

(Progeny Male &
Female ♂♂ & ♀
♂ ♂ ♂ ♂ ♂)